

Kindly amend claims 40 and 49, as follows:

40. (Twice Amended) A DNA encoding a polypeptide product consisting of an amino acid sequence selected (according to the numbering as presented in Figure 2) from the group consisting of:

- (a) amino acids -26 through 194; ✓
- (b) amino acids 1 through 194; ✓
- [(c) a polypeptide of subpart (a) or (b) wherein one or more cysteine residues is deleted or replaced by alanine or serine and wherein said cysteine residues are selected from the group consisting of amino acid positions 1, 13, 72, 101, 126, 128, 133, 138, 146, 167 and 175 in Figure 2;
- 1 (d) a polypeptide of subpart (a) or (b) wherein one or more tyrosine residues is replaced by phenylalanine and wherein said tyrosine residues are selected from the group consisting of amino acid positions 36, 45, 64, 84, 122, 139 and 178 in Figure 2;
- (e) a polypeptide of any of subparts (a), (b), (c) or (d), lacking residues -26 through -1, and having a methionyl residue at position -1];
- (c) a polypeptide of subpart (a), lacking residues -26 through -1 and having a methionyl residue at position -1; and
- (d) a polypeptide of subpart (b), having a methionyl residue at position -1.

49. (Twice Amended) A process for the production of a polypeptide selected (according to the numbering as presented in Figure 2) from the group consisting of:

- (a) amino acids -26 through 194;
- (b) amino acids 1 through 194;
- (c) a polypeptide of subpart (a) or (b) wherein one or more cysteine residues is deleted or replaced by alanine or serine and wherein said cysteine residues are selected from the group consisting of amino acid positions 1, 13, 72, 101, 126, 128, 133, 138, 146, 167 and 175 in Figure 2;
- (d) a polypeptide of subpart (a) or (b) wherein one or more tyrosine residues is replaced by phenylalanine and wherein said tyrosine residues are selected from the group consisting of amino acid positions 36, 45, 64, 84, 122, 139 and 178 in Figure 2;
- (e) a polypeptide of any of subparts (a), (b), (c) or (d), lacking residues -26 through -1, and having a methionyl residue at position -1;

(c) a polypeptide of subpart (a), lacking residues -26 through -1 and having a methionyl residue at position -1; and

(d) a polypeptide of subpart (b), having a methionyl residue at position -1;

said process comprising:

growing, under suitable nutrient conditions, prokaryotic or eucaryotic host cells transformed [or transfected] with a DNA